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W P E R L H

(TW)

Release 3.1A John F. Collins, BioComputing Research Unit.
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MPsrch_pp protein - protein database search, using Smith-Waterman algorithm

Run on: Mon Oct 2 09:33:11 2000; MasPar time 6.68 Seconds
Tabular output not generated. 436.063 Million cell updates/sec

Title: >US-09-381-497-2
Description: (1-123) from US09381497.pep
Perfect Score: 903
Sequence: 1 EVLVESGGGLVKPGGSLKL.....SSYGVLFAWGQGLTVTUSA 123

Scoring table: PAM 150
Gap 11

Searched: 188963 seqs, 23686106 residues

Post-processing: Minimum Match 0%
Listing first 45 summaries

Database: a-geneseq36
1:geneseqp

Statistics: Mean 30.324; Variance 152.652; scale 0.199

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query Match	Score	Length	ID	Description	Pred. No.
1	896	99.2	123	1	W66099 anti-CD22 monoclonal a	1.58e-63
2	748	82.8	121	1	W86125 protein sequence of mo	3.76e-51
3	744	82.4	121	1	W22951 Monoclonal antibody (M	8.11e-51
4	744	82.4	121	1	W86118 Murine 340 VH amino ac	8.11e-51
5	735	81.4	139	1	W21656 Chimeric MAB 15 PCR-mo	4.56e-50
6	730	80.8	140	1	W21654 Mouse MAB 15 heavy cha	1.19e-49
7	728	80.6	136	1	R06251 Variable region of mur	1.75e-49
8	717	79.4	139	1	R52773 Murine KC-4 immunoglob	1.44e-48
9	717	79.4	139	1	R52791 Murine KC-4 immunoglob	1.44e-48
10	716	79.3	118	1	W89627 Mouse humanised antibo	1.74e-48
11	716	79.3	118	1	W57576 Chimeric H chain SEQ I	1.74e-48
12	716	79.3	137	1	W89625 Mouse humanised antibo	1.74e-48
13	716	79.3	137	1	W57592 Chimeric antibody agai	1.74e-48
14	716	79.3	247	1	W11917 Murine MAB SK48-E26 he	1.74e-48
15	713	79.0	121	1	W86122 Protein sequence of de	3.10e-48
16	713	79.0	123	1	W08582 Human antibody C4.1 he	3.10e-48
17	712	78.8	121	1	W86120 Protein sequence of hu	3.75e-48
18	712	78.8	139	1	W21652 Humanised reshaped MAB	3.75e-48
19	710	78.6	138	1	R20064 MRK16-H chain.	5.50e-48
20	709	78.5	117	1	W76003 LM609 antibody heavy c	6.67e-48
21	706	78.2	142	1	R30882 Antibody 4A2 heavy cha	1.18e-47
22	704	78.0	138	1	W03722 Anti-human gp39 MAB 39	1.74e-47
23	703	77.9	123	1	R43827 Anti-lysozyme VH.	2.10e-47

24	703	77.9	123	1	R45187 Heavy chain variable d	2.10e-47
25	700	77.5	119	1	W73503 Antibody 15D3 heavy ch	3.74e-47
26	700	77.5	119	1	W69322 15D3 antibody heavy ch	3.74e-47
27	700	77.5	158	1	W19577 Mouse anti-idiotypic a	3.74e-47
28	700	77.5	158	1	W19579 Mouse anti-idiotypic a	3.74e-47
29	698	77.3	119	1	W29996 Humanised variant of h	5.48e-47
30	697	77.2	120	1	W00240 EGF receptor chimeric	6.64e-47
31	696	77.1	117	1	R79155 Human Ige receptor-bln	8.04e-47
32	696	77.1	117	1	W27354 Heavy chain variable r	8.04e-47
33	696	77.1	117	1	W27357 Heavy chain variable r	8.04e-47
34	693	76.7	121	1	W86124 Protein sequence of al	1.43e-46
35	692	76.6	118	1	R41233 Monoclonal antibody BW	1.73e-46
36	692	76.6	119	1	R56239 VH region of anti-fuco	1.73e-46
37	692	76.6	138	1	R32246 BR55-2 murine IgG3 hea	1.73e-46
38	692	76.6	138	1	R32242 Chimeric MAB heavy cha	1.73e-46
39	692	76.6	139	1	R31588 BR55-2 heavy chain var	1.73e-46
40	691	76.5	119	1	W29994 Heavy chain variable r	2.09e-46
41	691	76.5	140	1	W06205 Xenograft antibody HAR	2.09e-46
42	689	76.3	448	1	R97376 Murine anti-BGH MAB he	3.07e-46
43	687	76.1	448	1	R43673 Mouse anti-bovine grow	4.50e-46
44	686	76.0	448	1	R06476 Heavy chain of anti-bo	5.45e-46
45	685	75.9	117	1	R79157 Human Ige receptor-bln	6.60e-46

ALIGNMENTS

RESULT	ID	W66099 standard; Protein; 123 AA.	W66099;	DT	10-DEC-1998 (first entry)	DE	anti-CD22 monoclonal antibody heavy chain variable region.	DW	anti-CD22 monoclonal antibody heavy chain variable region; VL;	KW	Pseudomonas exotoxin; variable heavy chain; VH; variable light chain;	OS	Mammalia	FH	Key	Location/Qualifiers	FT	Misc_difference 121	FT	Misc_difference 121	PN	WO9841641-A1.	PD	24-SEP-1998.	PF	19-MAR-1998; U05453	PR	20-MAR-1997; US-041437.	PI	(USSH) US DEPT HEALTH & HUMAN SERVICES.	PA	Fitzgerald D, Kraitman R, Mansfield E, Pastan I;	DR	WPI: 98-521227/44.	DR	N-PSDB: V07642.	PT	Recombinant anti-CD22 antibodies and immuno-conjugates - of	PT	antibodies linked to a therapeutic agent, e.g. Pseudomonas exotoxin	PT	or a label; for inhibiting malignant B-cells	PS	Claim 6; Fig 1; 71pp; English.	CC	The invention claims for a recombinant immunoconjugate comprising	CC	of a therapeutic agent (e.g. Pseudomonas exotoxin) or a detectable	CC	label peptide bonded to a recombinant anti-CD22 antibody (RFB4 IgG)	CC	having the present variable heavy (VH) chain with a cysteine residue	CC	at amino acid 44 and a variable light (VL; W66098) chain with a	CC	cysteine residue at amino acid 100. The immunoconjugate is claimed	CC	to inhibit the growth of malignant B-cells in vivo, such as rodent,	CC	canine or primate B-cells. The anti-CD22 antibody is claimed useful	CC	for detecting CD22 protein in a sample or in vivo in a mammal, and	CC	can be used in diagnostic kits.	SC	Sequence 123 AA;
--------	----	-----------------------------------	---------	----	---------------------------	----	--	----	--	----	---	----	----------	----	-----	---------------------	----	---------------------	----	---------------------	----	---------------	----	--------------	----	---------------------	----	-------------------------	----	---	----	--	----	--------------------	----	-----------------	----	---	----	---	----	--	----	--------------------------------	----	---	----	--	----	---	----	--	----	---	----	--	----	---	----	---	----	--	----	---------------------------------	----	------------------

Query Match	99.2%;	Score 896;	DB 1;	Length 123;
Best Local Similarity	99.2%;	Pred. No. 1.58e-63;		
Matches	122;	Conservative	0;	Mismatches 1;
Indels	0;	Gaps	0;	
Db	1	EVLVESGGGLVKPGGSLKLSCAASGFAFSIYDMSVWROTPEKRLWEWAYISSGGGTTY 60		
Qy	1	EVLVESGGGLVKPGGSLKLSCAASGFAFSIYDMSVWROTPEKRLWEWAYISSGGGTTY 60		
Db	61	PDTVKGRFTISRDNAKNTLYLQMSLKSEDTAMYYCARHSGYSSGYVLFAWGQGLTVT 120		
Qy	61	PDTVKGRFTISRDNAKNTLYLQMSLKSEDTAMYYCARHSGYSSGYVLFAWGQGLTVT 120		

PD 26-NOV-1998.
 PR 21-MAY-1998; G01473.
 PR 14-APR-1998; GB-007751.
 PR 21-MAY-1997; GB-010480.
 PR 31-JUL-1997; GB-016197.
 PR 28-NOV-1997; GB-025270.
 PR 02-DEC-1997; US-067235.
 PA (BIOV-) BIOVATION LTD.
 PI Carr FJ;
 DR WPI; 99-045301/04.
 DR N-PSDB; V81002.
 PT Reducing immunogenicity of proteins - by modifying the amino acid
 PT sequence of the protein to eliminate potential epitopes for T-cells
 PT of a given species
 PS Example 1; Fig 2; 77pp; English.
 CC The invention relates to a method for the production of non-immunogenic
 CC proteins. The method comprises determining at least part of the amino
 CC acid sequence of the protein; (b) identifying in the amino acid sequence
 CC one or more potential epitopes for T-cells (T-cell epitopes) of the given
 CC species; and (c) modifying the amino acid sequence to eliminate at least
 CC one of the T-cell epitopes identified in step (b) thereby to eliminate or
 CC reduce the immunogenicity of the protein when exposed to the immune
 CC system-of-the-given-species. A method of analysing a pre-existing protein
 CC to predict the basis for immunogenic responses is also provided. The
 CC methods can be used particularly for reducing the immunogenicity of
 CC immunoglobulins or therapeutic proteins, e.g. Streptokinase (SK). The
 CC products can be used for diagnosis and therapy. The present sequence
 CC represents the amino acid sequence of murine 340 Vh.
 SQ Sequence 121 AA;

Query Match 82.4%; Score 744; DB 1; Length 121;
 Best Local Similarity 83.7%; Pred. No. 8,11e-51;
 Matches 103; Conservative 7; Mismatches 11; Indels 2; Gaps 1;

Db 1 EVQLVESGGGLVKGAGSLKLSCAASGFAFDYDMSWVROTPEKRLWVAYIGSGGRTYY 60
 QY 1 EVQLVESGGGLVKGAGSLKLSCAASGFAFDYDMSWVROTPEKRLWVAYISGGGTTY 60

Db 61 PDTVKGRFTISRDNKNTLYLQNSLKSEDTAMYYCARHGHVYDVAV--DYWGQGTSTV 118
 QY 61 PDTVKGRFTISRDNKNTLYLQNSLKSEDTAMYYCARHGHVYDVAV--DYWGQGTSTV 118

Db 119 VSS 121
 QY 121 VSA 123

RESULT 5
 ID W21656 standard; Protein; 139 AA.

AC W21656;
 DT 03-JAN-1998 (first entry)
 DE Chimeric MAB 15 PCR-modified heavy chain variable region.
 KW Humanised antibody; monoclonal antibody; MAB 15; tumour;
 KW lung cancer; therapy.
 OS Chimeric Mus musculus.
 OS Chimeric synthetic.

FT Key Location/Qualifiers
 FT Peptide 1..19
 FT /label= Sig_peptide
 FT Protein 25..139
 FT /label= Mat_protein
 FT Region 20..49
 FT /label= Framework-1
 FT Region 50..54
 FT /label= CDR1
 FT /note= "complementarity determining region 1"
 FT Region 55..68
 FT /label= Framework-2
 FT Region 69..86
 FT /label= CDR2
 FT /note= "complementarity determining region 2"
 FT Region 87..117
 FT /label= Framework-3
 FT /note= "complementarity determining region 3"

FT Region 118..128
 FT /label= CDR3
 FT /note= "complementarity determining region 3"
 FT 129..138
 FT /label= Framework-4
 PN EP-781847-A1.
 PD 02-JUL-1997.
 PD 25-OCT-1996; 117154.
 PR 06-NOV-1995; EP-117407.
 PR (MERE) MERCK PATENT GMBH.
 PI Bendig M, Jones T, Saldana J;
 DR WPI; 97-334904/31.
 DR N-PSDB; T72269.
 PT Humanised form of murine monoclonal antibody MAB 15 - useful for
 PT treating lung cancer
 PS Disclosure; Fig 5; 71pp; English.
 CC This polypeptide comprises the heavy chain variable region VH
 CC region of murine monoclonal antibody (MAB) 15 (DSM ACC2117). It is
 CC encoded by a MAB 15 VH cDNA sequence (72269) modified for the
 CC expression of chimeric antibody. The VL sequence was similarly
 CC obtained (see W21655). The modified VH and VL sequences were used
 CC in a claimed process to model and design novel humanised, reshaped
 CC MAB 15 having humanised, reshaped VH and VL sequences (see W21652
 CC and W21651), which can be used for treating tumours, especially
 CC lung cancer, and for the manufacture of a drug related to tumours,
 CC especially lung cancer.
 SQ Sequence 139 AA;

Query Match 81.4%; Score 735; DB 1; Length 139;
 Best Local Similarity 82.9%; Pred. No. 4.56e-50;
 Matches 102; Conservative 9; Mismatches 9; Indels 3; Gaps 1;

Db 20 EVQVSEGGGLVKGAGSLKLSCAASGFAFDYDMSWVROTPEKRLWVAYLSRGGSTYY 79
 QY 1 EVQVSEGGGLVKGAGSLKLSCAASGFAFDYDMSWVROTPEKRLWVAYISGGGTTY 60

Db 80 PDTVKGRFTISRDNKNTLYLQNSLKSEDTAMYYCARHGHVYDVAV--FDYWGQGTSTV 136
 QY 61 PDTVKGRFTISRDNKNTLYLQNSLKSEDTAMYYCARHGHVYDVAV--FDYWGQGTSTV 120

Db 137 VSA 139
 QY 121 VSA 123

RESULT 6
 ID W21654 standard; Protein; 140 AA.

AC W21654;
 DT 03-JAN-1998 (first entry)
 DE Mouse MAB 15 heavy chain variable region.
 KW Humanised antibody; monoclonal antibody; MAB 15; tumour;
 KW lung cancer; therapy.
 OS Mus musculus.

FT Key Location/Qualifiers
 FT Peptide 1..19
 FT /label= Sig_peptide
 FT Protein 20..140
 FT /label= Mat_protein
 FT Region 20..49
 FT /label= Framework-1
 FT Region 50..53
 FT /label= CDR1
 FT /note= "complementarity determining region 1"
 FT Region 54..68
 FT /label= Framework-2
 FT Region 69..86
 FT /label= CDR2
 FT /note= "complementarity determining region 2"
 FT Region 87..117
 FT /label= Framework-3
 FT /label= CDR3
 FT /note= "complementarity determining region 3"

```
FT Region 129..139
FT /label= Framework-4
PN EP-781847-A1.
PD 02-JUL-1997.
PF 25-OCT-1996; 117154.
PR 06-NOV-1995; EP-117407.
PA (WERE ) MERCK PATENT GMBH.
PI Bendig M, Jones T, Saldana J;
DR N-PSDB; T72267.
DR Humanised form of murine monoclonal antibody MAB 15 - useful for
PT treating lung cancer
PT Example 1; Fig 2; 71pp; English.
PS This polypeptide comprises the heavy chain variable region VH
CC of murine monoclonal antibody (MAB) 15 (DSM ACC2117), a MAB that
CC shows a therapeutic effect on human tumour cells, especially human
CC lung cancer. Its sequence was deduced from an isolated cDNA
CC cloner (see T72267). The MAB 15 VL region sequence (W21653)
CC has also been determined. Amplified VH and VL cDNA sequences were
CC used in a claimed process for the production of novel humanised,
CC reshaped MAB 15 having humanised, reshaped VH and VL regions (see
CC W21652 and W21651), which can be used for treating tumours,
CC especially lung cancer, and for the manufacture of a drug related
CC to tumours, especially lung cancer.
SQ Sequence 140 AA;

Query Match 80.8%; Score 730; DB 1; Length 140;
Best Local Similarity 82.1%; Pred. No. 1.19e-49;
Matches 101; Conservative 10; Mismatches 9; Indels 3; Gaps 1;

Db 20 EVQVSEGGGLVPGGSLKLSCAASGFAFSDYDMSWVRQTPKRLWVAYISGGGTTY 79
QY 1 EVQVSEGGGLVPGGSLKLSCAASGFAFSDYDMSWVRQTPKRLWVAYISGGGTTY 60

Db 80 PDTVKGRFTISRDNKAKILFLQWTSLSKSEDAAMYICARHGEVPRW---FDYWGQGLT 136
QY 61 PDTVKGRFTISRDNKAKNTLYLQMSLSKSEDAAMYICARHSGYSGYGLFAYWGQGLT 120

Db 137 VSA 139
QY 121 VSA 123

RESULT 7
ID R06251; standard; protein; 136 AA.
AC R06251;
DT 10-DEC-1990---(first-entry)
DE Variable region of murine AHT 54 heavy chain.
KW Interleukin-2 receptor; IL-2; tumour necrosis factor; TNF; ss.
OS Mus sp.
PN EP-380068-A.
PD 01-AUG-1990.
PF 24-JAN-1990; 101351.
PR 24-JAN-1989; US-301216.
PR 04-DEC-1989; US-441702.
PA (MOLE-) MOLECULAR THERAPU.
PI Zarler B;
DR WPI; 90-232892/31.
DR N-PSDB; Q05555.
PT Expression vectors for producing chimeric monoclonal antibodies -
PT which express human constant region and non-human variable region
PS Disclosure; p; English.
CC Mabs comprising mouse CH and CL constant regions which human
CC variable regions may be used to create mouse/human hybrid Mabs,
CC which have a longer serum half-life. Method can be used to produce
CC Abs against interleukin-2 receptor and tumour necrosis factor.
SQ Sequence 136 AA;

Query Match 80.6%; Score 728; DB 1; Length 136;
Best Local Similarity 88.6%; Pred. No. 1.75e-49;
Matches 109; Conservative 3; Mismatches 5; Indels 6; Gaps 1;

Db 20 EVQVSEGGGLVPGGSLKLSCAASGFAFSSIDMSWVRQTPKRLWVAYISGGGNTYY 79
```

```
QY 1 EVQVSEGGGLVPGGSLKLSCAASGFAFSDYDMSWVRQTPKRLWVAYISGGGTTY 60
Db 80 PDTVKGRFTISRDNKAKNTLYLQMSLSKSEDAAMYICARR-----YGLPFAYWGQGLT 133
QY 61 PDTVKGRFTISRDNKAKNTLYLQMSLSKSEDAAMYICARHSGYSGYGLFAYWGQGLT 120
Db 134 VSA 136
QY 121 VSA 123

RESULT 8
ID R52773; standard; Protein; 139 AA.
AC R52773;
DT 24-JAN-1995 (first entry)
DE Murine KC-4 immunoglobulin heavy chain variable region (deduced).
KW Immunoglobulin variable domain; primer; polymerase chain reaction;
KW chimeric antibody; human milk fat globule; human breast carcinoma;
KW murine anti-human carcinoma monoclonal antibody KC-4.
OS Mus musculus.
FH Key Location/Qualifiers
FT Protein 20..139
FT /label= KC-4_mature_VL-chain
FT region 20..49
FT /label= FR1
FT region 50..54
FT /label= CDR1
FT region 55..68
FT /label= FR2
FT region 69..85
FT /label= CDR2
FT region 86..117
FT /label= FR3
FT region 118..128
FT /label= CDR3
FT region 129..139
FT /label= FR4
PN W09411508-A.
PD 26-MAY-1994.
PF 15-NOV-1993; U11316.
PR 13-NOV-1992; US-977706.
PR 13-NOV-1992; US-977707.
PR 28-SEP-1993; US-128015.
PA (CANC-) CANCER RES FUND CONTRA COSTA.
DR WPI; 94-183509/22.
DR N-PSDB; -Q62764-
PT Chimeric human-murine polypeptide(s) specific for human mammary
PT fat globule antigen - for imaging, diagnosing and treating
PT neoplasia, with less undesirable immunogenic response
PS Example 27; Page 41; 54pp; English.
CC An initial isolation of cDNAs coding for murine anti-human breast
CC carcinoma MAB KC-4 was performed using PCR with commercially
CC available primers (see Q62751-Q62758, available from NOVAGEN).
CC Subsequent cloning using PCR primers JO20, JO21, JO22 and JO24
CC (see Q62759-Q62762) resulted in the isolation of the mouse Ig
CC variable domains. The amplified cDNAs were sequenced (Q62763 and
CC Q62764) and amino acid sequences were deduced from them. Chimeric
CC mouse-human antibodies were constructed using human constant
CC regions so as to produce less immunogenic polypeptides which
CC retained the anti-human carcinoma binding specificity of KC-4.
SQ Sequence 139 AA;

Query Match 79.4%; Score 717; DB 1; Length 139;
Best Local Similarity 82.1%; Pred. No. 1.44e-48;
Matches 101; Conservative 8; Mismatches 11; Indels 3; Gaps 3;

Db 20 EVQVSEGGGLVPGGSLKLSCAASGFAFSSYAMSVROSPEKRLWVAEISSGGNYAY 79
QY 1 EVQVSEGGGLVPGGSLKLSCAASGFAFSDYDMSWVRQTPKRLWVAYISGGGTTY 60
Db 80 QDTVKGRFTISRDNKAKNTLYLQMSLSKSEDAAMYICAREY---YGLPFAYWGQGLT 136
QY 121 VSA 123
```

OY 61 PTVKGRFTISRDNKNTLYLQMSLKSEDTAMFYCARHSGYSSYGVLFAYWGQGLTV 120
 DB 137 VSA 139
 OY 121 VSA 123

RESULT 9

ID R52791 standard; Protein: 139 AA.

AC R52791:

DT 24-JAN-1995 (first entry)

DE Murine KC-4 immunoglobulin heavy chain variable region (deduced).

KM Immunoglobulin variable domain; primer: polymerase chain reaction;

KW chimeric antibody; human milk fat globule; human breast carcinoma;

OS murine anti-human carcinoma monoclonal antibody KC-4.

PI Mus musculus.

PH Key Location/Qualifiers

FT protein 20..139

FT region /label- KC-4_mature_VL-chain

FT region 20..49

FT region /label- FR1

FT region 50..54

FT region /label- CDRI

FT region 55..68

FT region /label- FR2

FT region 69..85

FT region /label- CDR2

FT region 86..117

FT region /label- FR3

FT region 118..128

FT region /label- CDR3

FT region 129..139

FT region /label- FR4

PN WO9411509-A.

PD 26-MAY-1994.

PE 16-NOV-1993: U11445.

PF 16-NOV-1993: US-977696.

PR 30-SEP-1993: US-129930.

PR 08-OCT-1993: US-134346.

PA (CANC-) CANCER RES FUND CONTRA COSTA.

PI Certian RL, Do Couto FJR, Padlan EA, Peterson JA;

DR WPI: 94-183510/22.

PT New analogue peptide(s) comprising antibody variable regions -

PT used to develop prods. for use in the detection, diagnosis,

PT therapy and prevention of neoplasms

PS Example 26; Page 61; 109pp; English.

CC An initial isolation of cDNAs coding for murine anti-human breast

CC carcinoma MAb KC-4 was performed using PCR with commercially

CC available primers (see 062776-062783, available from NOVAGEN).

CC Subsequent cloning using PCR primers JO20, JO21, JO22 and JO24

CC (see 062784-062787) resulted in the isolation of the mouse Ig

CC variable domains. The amplified cDNAs were sequenced (062788 and

CC 062789) and amino acid sequences were deduced from them. Chimeric

CC mouse-human antibodies were constructed using human constant

CC regions so as to produce less immunogenic polypeptides which

CC retained the anti-human carcinoma binding specificity of KC-4.

CC Sequence 139 AA;

Query Match 79.4%; Score 717; DB 1; Length 139;

Best Local Similarity 82.1%; Pred. No. 1.44e-48;

Matches 101; Conservative 8; Mismatches 11; Indels 3; Gaps 3;

DB 20 EVQWVESGGGLVPGGSLKSCAASGFAFSYAMSVWRQSPERLEWVAEISSGGYAYY 79

OY 1 EVQWVESGGGLVPGGSLKSCAASGFAFSYIDMSVWRQTPERLEWVAEISSGGYAYY 60

DB 80 ODVTGRTFTISRDNKNTLYLEMSLSRSDTAMFYTCARD-YGIP-A-WFAYWGQGLTVS 136

OY 61 PTVKGRFTISRDNKNTLYLQMSLKSEDTAMFYCARHSGYSSYGVLFAYWGQGLTV 120

DB 137 VSA 139

OY 121 VSA 123

RESULT 10
 ID W89627 standard; Protein: 118 AA.

AC W89627:

DT 14-APR-1999 (first entry)

DE Mouse humanised antibody #23-57-137-1 heavy chain mature protein.

KM Human; parathyroid hormone related protein; PTHrP; cachexia; cancer;

KW Inhibitor; humanised.

OS Mus sp.

OS Synthetic.

PN WO9851329-A1.

PD 19-NOV-1998.

PE 13-MAY-1998; J02116.

PR 18-JUL-1997; JP-194445.

PR 15-MAY-1997; JP-125505.

PA (CHUS) CHUGAI SEIYAKU KK.

PI Ishii K, Sato K, Tuenari T;

DR WPI: 99-070101/06.

PT Inhibitors of binding of parathyroid hormone related peptide to its

PT receptor - useful for, e.g. treatment of cachexia arising from

PT cancer or other diseases

PS Example 2; Page 72-73; 125pp; Japanese.

CC The present invention describes compositions for the treatment of

CC cachexia containing a substance which inhibits the binding of a

CC parathyroid hormone related peptide (PTHrP) to its receptor, as an

CC active component. This substance may be an antagonist to the receptor,

CC or an antibody (preferably monoclonal) or antibody fragment,

CC recognising PTHrP. The antibody is preferably humanised or chimeric.

CC The present invention also describes a humanised antibody prepared

CC by hybridoma 23-57-137-1 (FERM BP-5631). The composition is used for

CC the treatment of cachexia arising in connection with diseases such as

CC cancer, thereby improving the quality of life of the patient. The

CC present sequence represents mouse humanised antibody heavy chain from

CC #23-57-137-1 from the present invention.

CC Sequence 118 AA;

Query Match 79.3%; Score 716; DB 1; Length 118;

Best Local Similarity 82.9%; Pred. No. 1.74e-48;

Matches 102; Conservative 9; Mismatches 7; Indels 5; Gaps 2;

DB 1 EVQWVESGGGLVPGGSLKSCAASGFTFSSYMSMTIRQTPDKRLWVAEISSGGSTYY 60

OY 1 EVQWVESGGGLVPGGSLKSCAASGFAFSYIDMSVWRQTPERLEWVAEISSGGSTYY 60

DB 61 PDVVKGRFTISRDNKNTLYLQMSLKSEDTAMFYCARQTT---MTY---FAYWGQGLTV 115

OY 61 PTVKGRFTISRDNKNTLYLQMSLKSEDTAMFYCARHSGYSSYGVLFAYWGQGLTV 120

DB 116 VSA 118

OY 121 VSA 123

RESULT 11
 ID W57576 standard; Protein: 118 AA.

AC W57576:

DT 03-SEP-1998 (first entry)

DE Chimeric H chain SEQ ID NO:46 for an antibody against hPTHrP.

KM Chimeric; antibody; human parathormone related peptide; hPTHrP; mouse;

KW L chain; H chain; hypercalcaemia; cancer; malignant lymphoma; CDR;

KW hypophosphemia; pathogen; vitamin D resistance; V region; C region;

KW humanised.

OS Synthetic.

OS Chimeric - Mus sp.

OS Chimeric - Homo sapiens.

PN WO9813388-A1.

PD 02-APR-1998.

PE 24-SEP-1997; J03382.

PR 24-JUL-1997; JP-214168.

PR 26-SEP-1996; JP-255196.

PA (CHUS) CHUGAI SEIYAKU KK.

PI Sato K, Wakahara Y, Yabuta N;

QY 121 VSA 123

RESULT 14

ID W11917 standard; Protein; 247 AA.
AC W11917;
DE 24-JUN-1997 (first entry)
DE Murine MAb SK48-E26 heavy chain.
KW Interleukin-1 beta; IL-1 beta; recombinant antibody;
KW humanised antibody; chimeric antibody; antibody engineering;
KW monoclonal antibody; MAB; SK48-E26; inflammation; therapy.
OS Homo sapiens.
FH Key Location/Qualifiers
FT peptide 1..19
FT /label= Sig_peptide
FT region 20..49
FT /label= FR1
FT /note= "framework region 1"
FT region 50..54
FT /label= CDR1
FT /note= "complementarity determining region 1
(Claim 10, page 48)"
FT region 55..68
FT /label= FR2
FT /note= "framework region 2"
FT region 69..85
FT /label= CDR2
FT /note= "complementarity determining region 2
(Claim 10, page 48)"
FT region 86..117
FT /label= FR3
FT /note= "framework region 3"
FT region 118..127
FT /label= CDR3
FT /note= "complementarity determining region 3
(Claim 10, page 48)"
FT region 128..138
FT /label= FR4
FT /note= "framework region 4"
FT region 139..247
FT /label= Constant_region
WO9501997-A1.
19-JAN-1995.
PF 07-JUL-1994; U07659.
PR 09-JUL-1993; US-090534.
PR 04-MAR-1994; US-206190.
PA (SMK) SMITHKLINE BEECHAM CORP.
PI Gross MS, Hurler MR, Jackson JR, Jonak ZL, Theisen TW;
PI Young PR;
PI WPI; 95-066868/09.
DR N-PSDB; T51436.
DR Recombinant and humanised chimeric antibodies against human
PT interleukin-1-beta - for preventing and treating
PT interleukin-mediated inflammatory disorders
PS Claim 5; Page 36-37; 62pp; English.
CC Amino acid sequences of the heavy chain (W11917) and light chain
CC (W11918) of anti-human interleukin-1 beta (IL-1 beta) murine
CC monoclonal antibody (MAB) SK48-E26 were deduced from nucleic acids
CC (T51436-37) derived from hybridoma SK48-E26. The heavy and light
CC chains, esp. the complementarity determining region sequences,
CC can be utilised in novel recombinant chimeric and humanised
CC antibodies (see also W11919-20) useful for the treatment and
CC prevention of IL-1 mediated inflammatory disorders.
SQ Sequence 247 AA;

Query Match 79.3%; Score 716; DB 1; Length 247;

Best Local Similarity 84.6%; Pred. No. 1.74e-48;

Matches 104; Conservative 5; Mismatches 10; Indels 4; Gaps 2;

Db 20 EVHVESGGGLVPGGSLKLSCAASGFTDTYDMSWVRQAPCKGLEWVAYISGGGTTY 79

QY 1 EVQLVESGGGLVPGGSLKLSCAASGFTDTYDMSWVRQAPCKGLEWVAYISGGGTTY 60

Db 80 PDTVKGRTISRDNKNTLYLQMSLSKSDTAMVYCARHSGYSSYGVLFAYWGQGLT 135
QY 61 PDTVKGRTISRDNKNTLYLQMSLSKSDTAMVYCARHSGYSSYGVLFAYWGQGLT 120

Db 136 VSS 138
QY 121 VSA 123

RESULT 15

ID W86122 standard; Protein; 121 AA.
AC W86122;
DE 03-MAR-1999 (first entry)
DE Protein sequence of de-immunised 340 Vh.
KW Non-immunogenic; epitope; T-cell; immunogenicity; immune system; SK;
KW immunoglobulin; therapeutic; streptokinase; de-immunised.
OS Homo sapiens.
PN WO9852976-A1.
PD 26-NOV-1998.
PF 21-MAY-1998; G01473.
PR 14-APR-1998; GB-007751.
PR 21-MAY-1997; GB-010480.
PR 31-JUL-1997; GB-016197.
PR 28-NOV-1997; GB-025270.
PR 02-DEC-1997; US-067235.
PA (BIOV-) BIOVATION LTD.
PI Carr FJ;
PI WPI; 99-045301/04.
PT Reducing immunogenicity of proteins - by modifying the amino acid
PT sequence of the protein to eliminate potential epitopes for T-cells
PT of a given species
PS Example 1; Fig 5; 77pp; English.
CC The invention relates to a method for the production of non-immunogenic
CC proteins. The method comprises determining at least part of the amino
CC acid sequence of the protein; (b) identifying in the amino acid sequence
CC one or more potential epitopes for T-cells (T-cell epitopes) of the given
CC species; and (c) modifying the amino acid sequence to eliminate at least
CC one of the T-cell epitopes identified in step (b) thereby to eliminate or
CC reduce the immunogenicity of the protein when exposed to the immune
CC system of the given species. A method of analysing a pre-existing protein
CC to predict the basis for immunogenic responses is also provided. The
CC methods can be used particularly for reducing the immunogenicity of
CC immunoglobulins or therapeutic proteins, e.g. Streptokinase (SK). The
CC products can be used for diagnosis and therapy. The present sequence
CC represents the protein sequence of de-immunised 340 Vh.
SQ Sequence 121 AA;

Query Match 79.0%; Score 713; DB 1; Length 121;

Best Local Similarity 78.9%; Pred. No. 3.10e-48;

Matches 97; Conservative 10; Mismatches 14; Indels 2; Gaps 1;

Db 1 EVQLVESGGGLVPGGSLKLSCAASGFTDTYDMSWVRQAPCKGLEWVAYISGGGTTY 60

QY 1 EVQLVESGGGLVPGGSLKLSCAASGFTDTYDMSWVRQAPCKGLEWVAYISGGGTTY 60

Db 61 PDTVKGRTISRDNKNTLYLQMSLSKSDTAMVYCARHSGYSSYGVLFAYWGQGLT 118

QY 61 PDTVKGRTISRDNKNTLYLQMSLSKSDTAMVYCARHSGYSSYGVLFAYWGQGLT 120

Db 119 VSS 121

QY 121 VSA 123

Search completed: Mon Oct 2 09:33:25 2000
Job time : 14 secs.

US-09-381-49

Query Match 97.0%; Score 734; DB 1; Length 131;
Best Local Similarity 97.2%; Pred. No. 9.32e-50;
Matches 104; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Db 21 DIQMTQTSSLSASLGDRVTISCRASQDISNYLNWYQOKPDGTVKLLIYYSRLHSGVPS 80
 Qy 1 DIQMTQTSSLSASLGDRVTISCRASQDISNYLNWYQOKPDGTVKLLIYYSILHSGVPS 60
 Db 81 RFSGSGSGTDYSLTISNLEQEDIATYFCQNGNTLPWTFGGGKTLEIK 127
 Qy 61 RFSGSGSGTDYSLTISNLEQEDFATYFCQNGNTLPWTFGGGKTLEIK 107